

In re Patent Application of

GABERT

Atty. Ref.: 1721-21

Serial No.

09/530.363

Group:

1656

Filed:

May 1, 2000

Examiner:

Spiegler

For:

METHOD FOR DIAGNOSING IN VITRO PATHOLOGIES

ASSOCIATED WITH GENE ARRANGEMENTS AND

DIAGNOSIS KITS

December 26, 2001

Assistant Commissioner for Patents Washington, DC 20231

Sir:

AMENDMENT UNDER RULE 116

Responsive to the Official Action dated September 25, 2001, entry and consideration of the following amendments and remarks are requested.

IN THE SPECIFICATION

Amend the specification as follows.

Page 21, delete the paragraph on line 16 and insert the following therefor:

--(SEQ ID NO 2, 44 nucleotides, Tm = 84°C).--

Page 21, delete the paragraph on line 26 and insert the following therefor:

--(SEQ ID NO 3, 32 nucleotides, Tm = 84°C), and--

Page 21, delete the paragraph on line 33 and insert the following therefor:

--(SEQ ID NO 4, 29 nucleotides, Tm = 86°C), and--

Page 22, delete the paragraph on line 33 and insert the following therefor:

AF

In re Patent Application of

GABERT

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Serial No. 09/530,363

Filed: May 1, 2000

Title: N

METHOD FOR DIAGNOSING IN VITRO PATHOLOGIES

ASSOCIATED WITH GENE ARRANGEMENTS AND DIAGNOSIS

KITS

Assistant Commissioner for Patents Washington, DC 20231

Sir:

RESPONSE UNDER RULE 116 EXPEDITED HANDLING PROCEDURES

Atty Dkt. 1721-21 C#/M#

Group Art Unit: 1656

Examiner: Spiegler

Date: December 26, 2001



RESPONSE/AMENDMENT/LETTER

This is a response/amendment/letter in the above-identified application and includes an attachment which is hereby incorporated by reference and the signature below serves as the signature to the attachment in the absence of any other signature thereon.

Fees are attached as calculated below:

Total effective claims after amendment 0 mir Previously paid for 20 (at least 20) = 0	nus highest number x \$ 18.00	\$	0.00
Independent claims after amendment 0 mir Previously paid for 3 (at least 3) = 0	nus highest number x \$ 84.00	\$	0.00
If proper multiple dependent claims now added for firs	t time, add \$280.00 (ignore improper)	\$	0.00
Petition is hereby made to extend the current due date so as to cover the filing date of this Paper and attachment(s) (\$110.00/1 month; \$400.00/2 months; \$920.00/3 months)		\$	0.00
Terminal disclaimer enclosed, add \$ 110.00		\$	0.00
First submission after Final Rejection pursuant to 37 C	CFR 1.129(a) (\$740.00)	\$	0.00
Second submission after Final Rejection pursuant to 3	7 CFR 1.129(a) (\$740.00)	\$	0.00
Please enter the previously unentered , filed			
	SUBTOTAL	\$	0.00
If "small entity," then enter half (1/2) of subtotal and su Statement filed herewith	btract	-\$	0.00
Rule 56 Information Disclosure Statement Filing Fee (\$180.00)	\$	0.00
Assignment Recording Fee (\$40.00)		\$	0.00
	TOTAL FEE ENCLOSED	\$	0.00

The Commissioner is hereby authorized to charge any <u>deficiency</u> in the fee(s) filed, or asserted to be filed, or which should have been filed herewith (or with any paper hereafter filed in this application by this firm) to our Account No. 14-1140. A duplicate copy of this sheet is attached.

1100 North Glebe Road, 8th Floor Arlington, Virginia 22201-4714 Telephone: (703) 816-4000 Facsimile: (703) 816-4100

BJS:1

NIXON & VANDERHYE P.C.

By Atty.: B. J. Sadoff, Reg. No. 36,663

Signature:

GABERT Serial No. 09/530,363

--(SEQ ID NO 5, 35 nucleotides, Tm = 81°C).--

Page 23, delete the paragraph spanning lines 5-6 and insert the following therefor:

--- ribozyme 1: <u>CUCCA</u>GCUGA UGAGUCCGUG AGGACGAAAC <u>CUUUGG</u> (SEQ ID NO 6)--

Page 23, delete the paragraph spanning lines 8-9 and insert the following therefor:

-- - ribozyme 2: <u>CUGGAAU</u>CUG AUGAGUCCGU GAGGACGAAA U<u>UUUCUUC</u> (SEQ ID NO 7)--

IN THE CLAIMS

Amend the claims as follows:

Cancel claim 16, without prejudice

- 17. (Amended) The method of claim 39, wherein the primers consist of 25 to 40 nucleotides.
- 18. (Amended) The method of claim 39, further comprising, labeling said PCR products, denaturing said labeled PCR products, and contacting the denatured labeled PCR products with a nucleotide sequence complementary to a fusion partner nucleotide sequence.